

[SEQ. ID NO: 3]

X-C-C-T-T-G-A-G-A-T-T-T-C-C-C-T-C
5'

G-G-A-A-C-T-C-T-A-A-A-G-G-G-A-G-X
[SEQ. ID NO: 4]

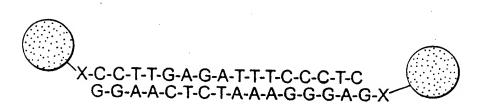


FIG. 2

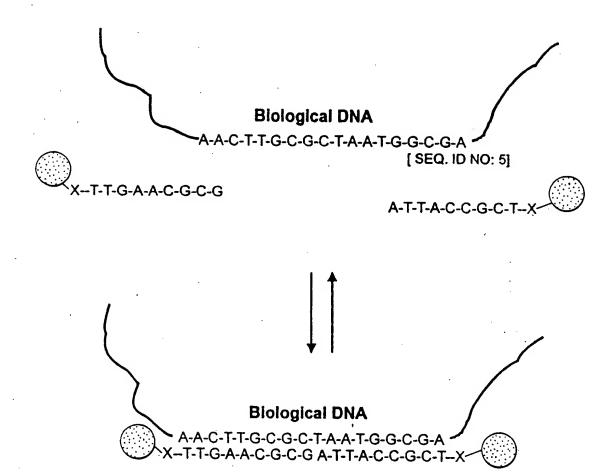
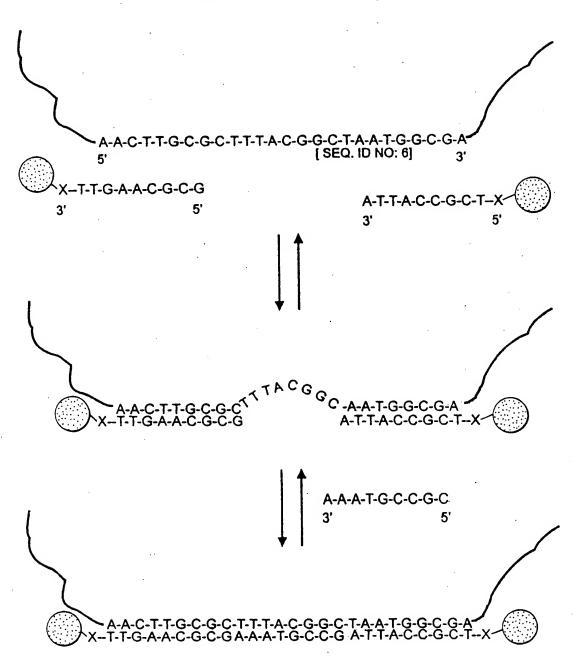
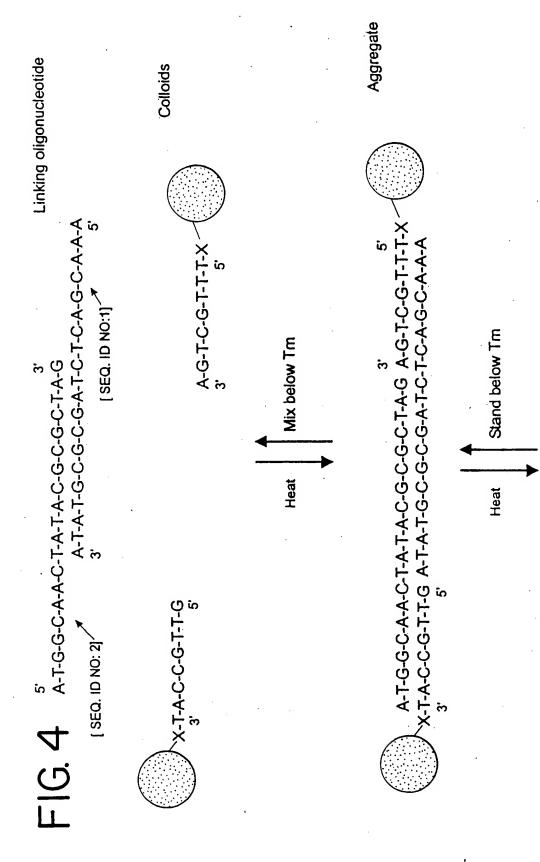


FIG. 3





Precipitate (formed by further cross-linking)

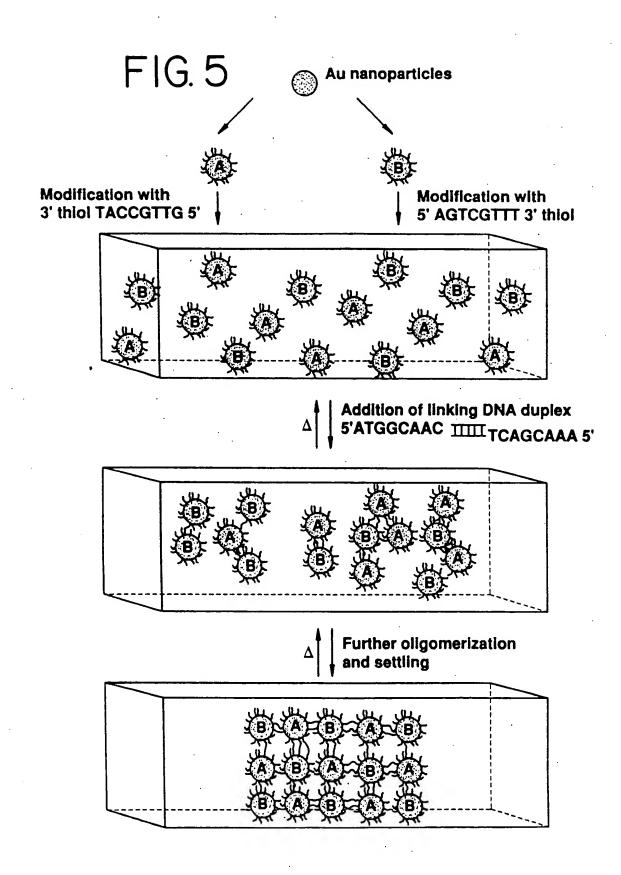
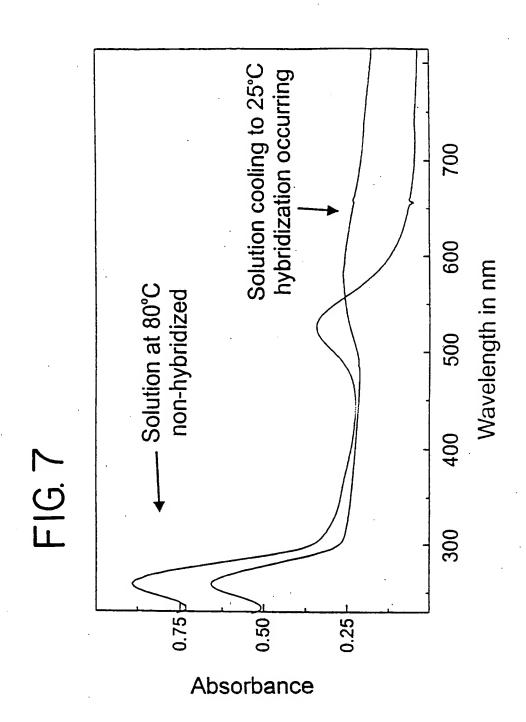


FIG. 6A FIG. 6B FIG. 6C









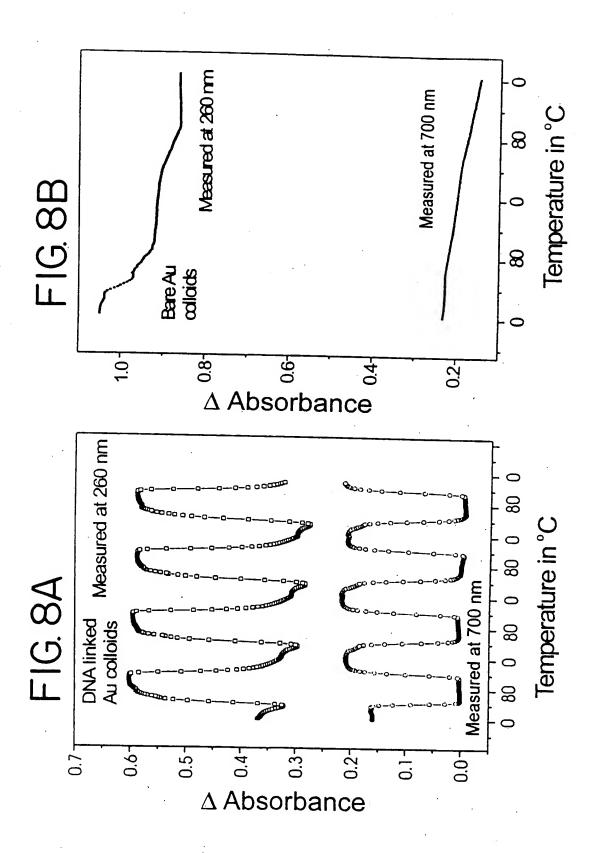


FIG. 9A

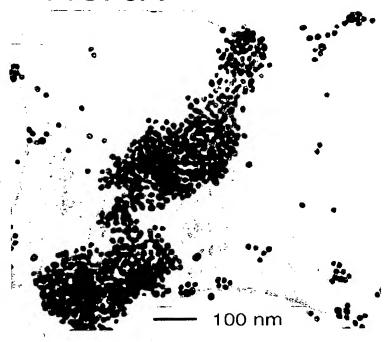
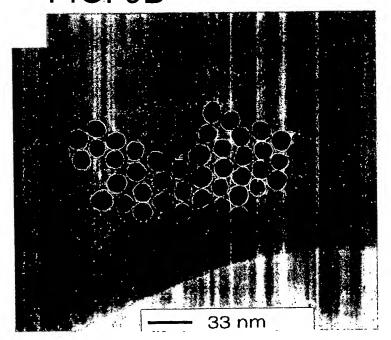
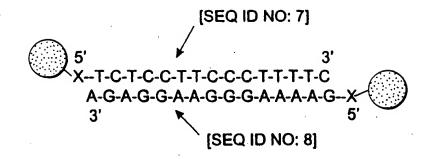
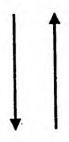
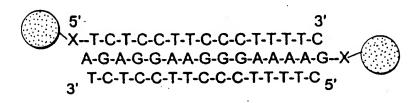


FIG. 9B









F1G. 1

`s-A_'T-G-G-C-A-A-C-T-A-T-A-C-G-C-G-C-T-A-G-A-G-T-C-G-T-T-T [SEQ. ID NO: 10]

T-A-C-C-G-T-T-G-A-T-A-T-G-C-G-C-G-A-T-C-T-C-A-G-C-A-A--S-C-3'
3'
[SEQ. ID NO: 11]

7-A-C-C-G-T-T-G-A-T-A-T-G-C-G-C-G-A-T-C-T-C-A-G-T-G-G-T-G-G-C-A-A-C-T-A-T-A-C-G-C-G-C-T-A-G-A-G-T-

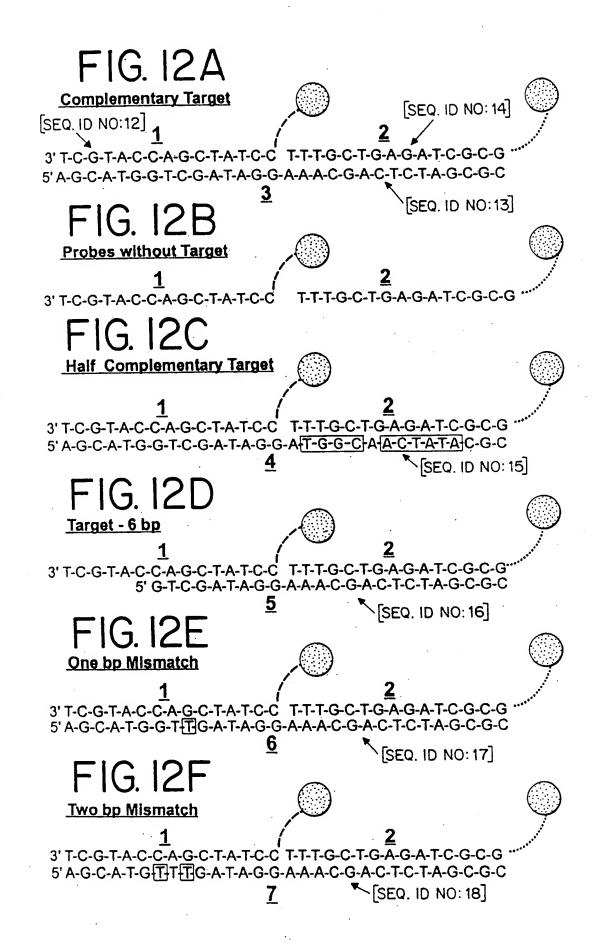
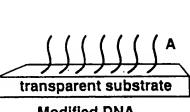
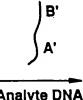


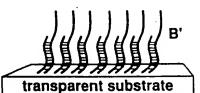
FIG. 13A



Modified DNA chemisorbed onto solid substrate

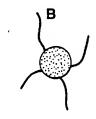


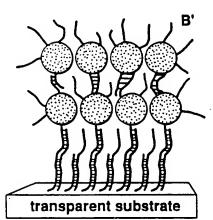
Analyte DNA



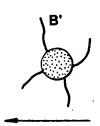
Analyte DNA hybridized onto substrate

DNA modified colloids





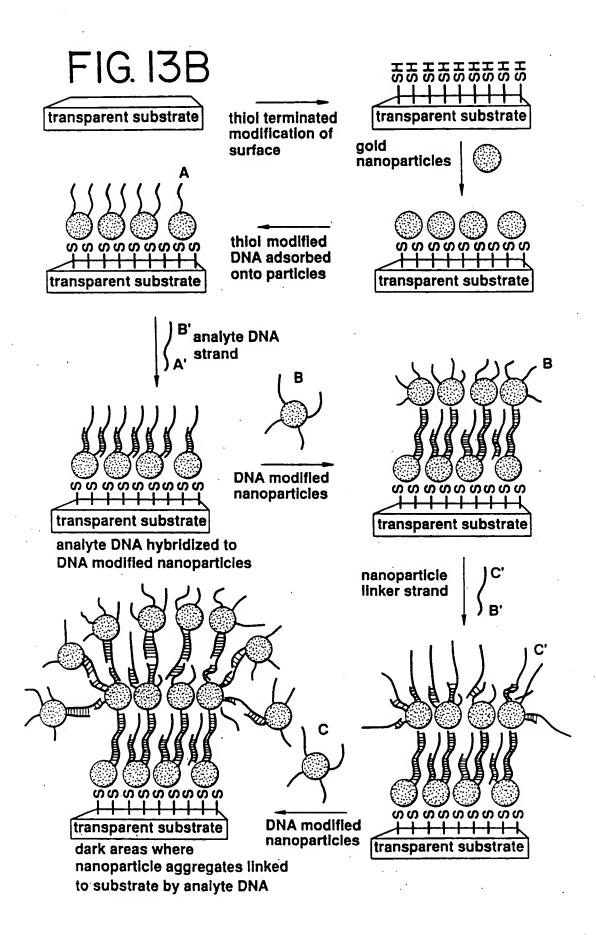
Dark areas where nanoparticle aggregates are linked to substrate surface by analyte DNA



transparent substrate

DNA modified colloids

hybridized to bound analyte DNA



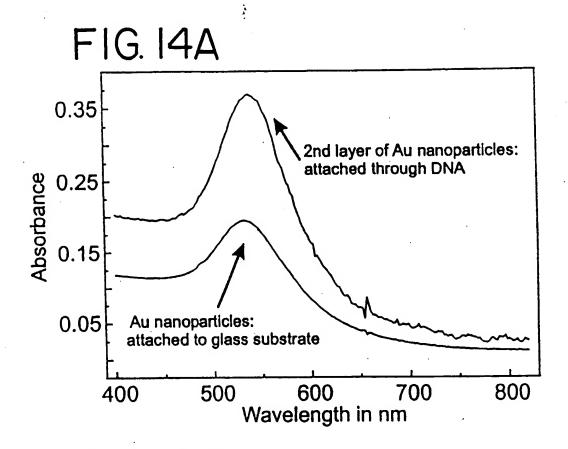


FIG. 14B

0.34

Eu 07

0.30

First Derivative: Tm=62 C

0.22

0 20 40 60 80

Temperature in °C

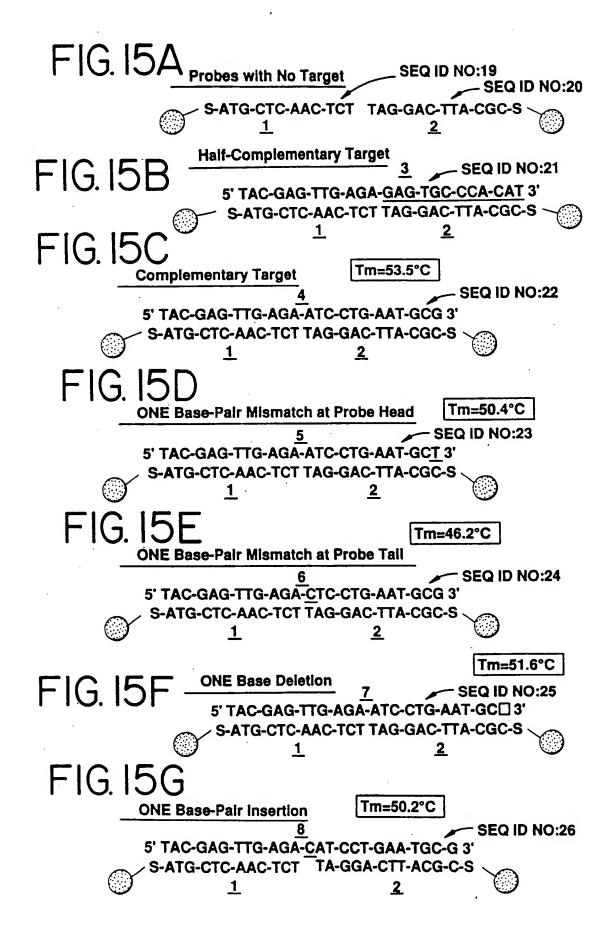


FIG. 16A

- S-ATG-CTC-AAC-TCT TAG-GAC-TTA-CGC-S 5' TAC-GAG-TTG-AGA-ATC-CTG-AAT-GCG 3'

24 Base Template

FIG. 16B

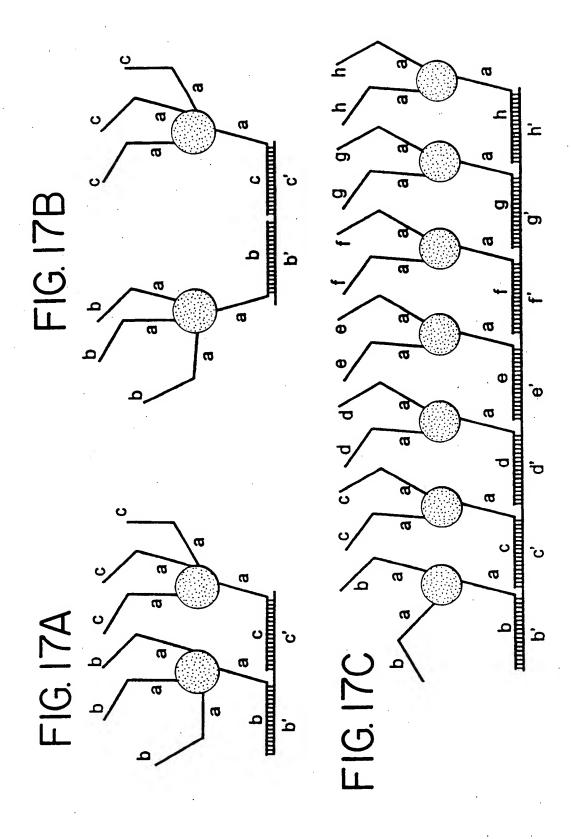
48 Base Template with Complementary 24 Base Filler

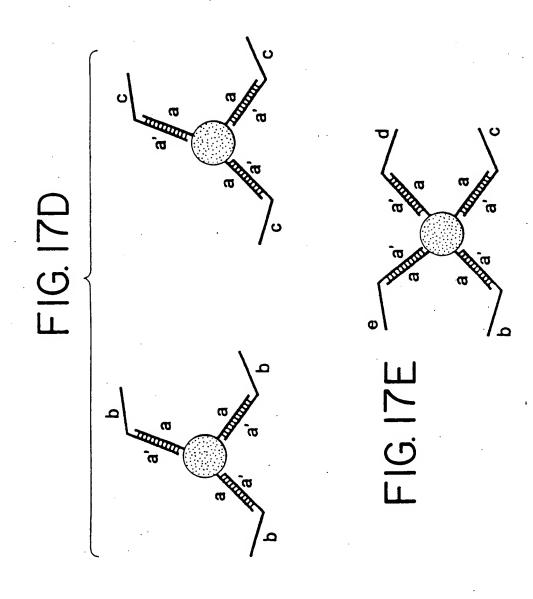
5' TAC-GAG-TTG-AGA-CCG-TTA-AGA-CGA-GGC-AAT-CAT-GCA-ATC-CTG-AAT-GCG 3' - S-ATG-CTC-AAC-TCT GGC-AAT-TCT-GCT-CCG-TTA-GTA-CGT TAG-GAC-TTA-CGC-S

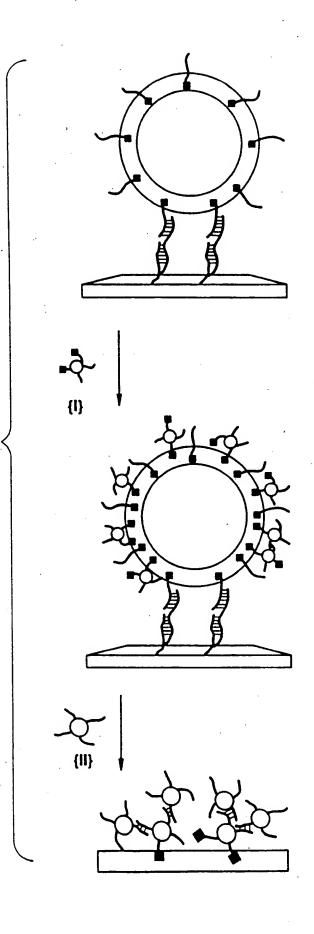
FIG. 16C

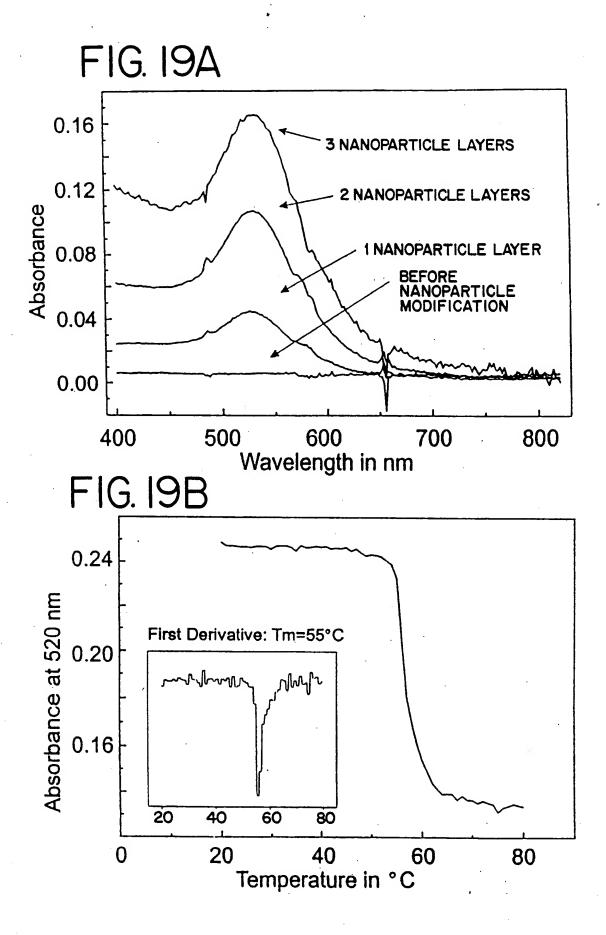
72 Base Template with Complementary 48 Base Filler

5' TAC-GAG-TTG-AGA-CCG-TTA-AGA-CGA-GGC-AAT-CAT-GCA-TAT-ATT-GGA-CGC-TTT-ACG-GAC-AAC-ATC-CTG-AAT-GCG 3'

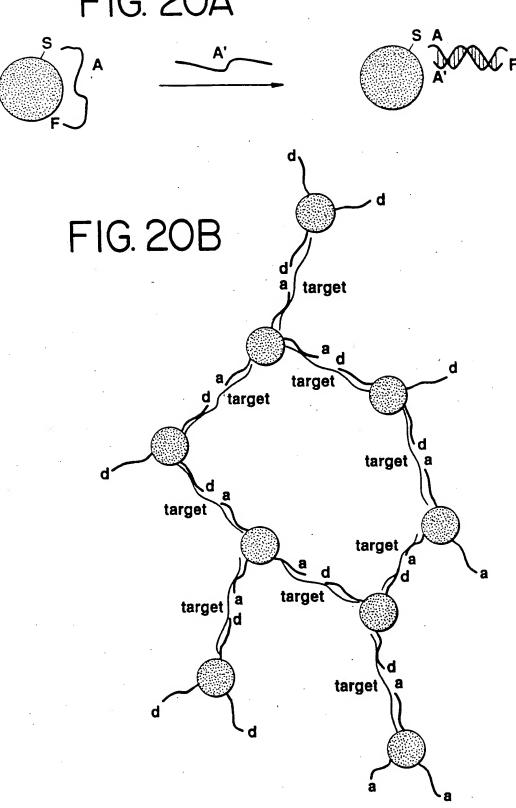


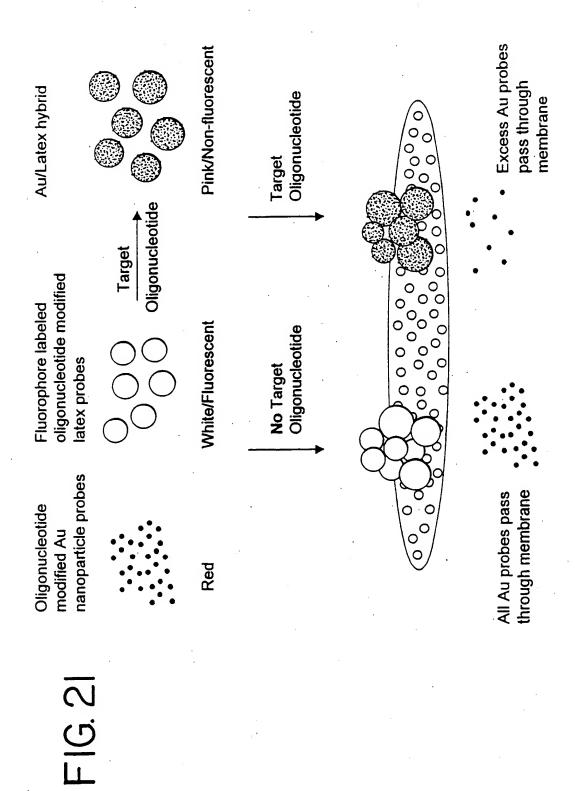


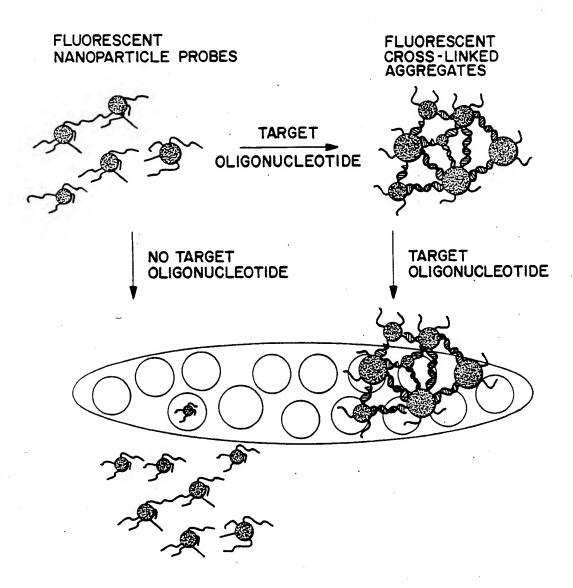












THE FLUORESCENT NANOPARTICLE PROBES PASS THROUGH THE MEMBRANE

THE FLUORESCENT CROSS-LINKED AGGREGATES ARE RETAINED BY THE MEMBRANE

Anthrax PCR Product

5'G GCG GAT GAG TCA GTA GTT AAG GAG GCT CAT AGA GAA GTA ATT AAT 3'C CGC CTA CTC AGT CAT CAA TTC CTC CGA GTA TCT CTT CAT TAA TTA

TCG TCA ACA GAG GGA TTA TTG TTA AAT ATT GAT AAG GAT ATA AGA AAA AGC AGT TGT CTC CCT AAT AAC AAT TTA TAA CTA TTC CTA TAT TCT TTT

ATA TTA TCC AGG GTT ATA TTG TAG AAA TTG AAG ATA CTG AAG GGC TT 3' TAT AAT AGG TCC CAA TAT AAC ATC TTT AAC TTC TAT GAC TTC CCG AA 5'

141 mer Anthrax PCR product [SEQ ID NO:36]

3' CTC CCT AAT AAC AAT

[SEQ ID NO:37]

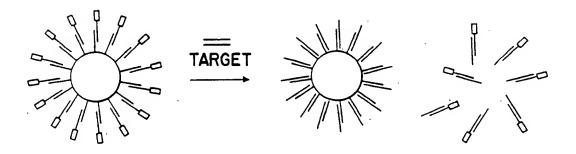
3' TTA TAA CTA TTC CTA -

[SEQ ID NO:38]

Oligonucleotide-Nanoparticle Probes

Blocker Oligonucleotides

3' C CGC CTA CTC AGT CAT CAA TTC CTC CGA GT	[SEQ ID NO:39]
3' A TCT CTT CAT TAA TTA AGC AGT TGT	[SEQ ID NO:40]
3' TAT TOT TTT TAT AAT AGG TCC CAA TAT	[SEQ ID NO:41]
3' AAC ATC TIT AAC TTC TAT GAC TTC CCG AA	[SEQ ID NO:42]



SATELLITE PROBE

DETECTION SIGNAL

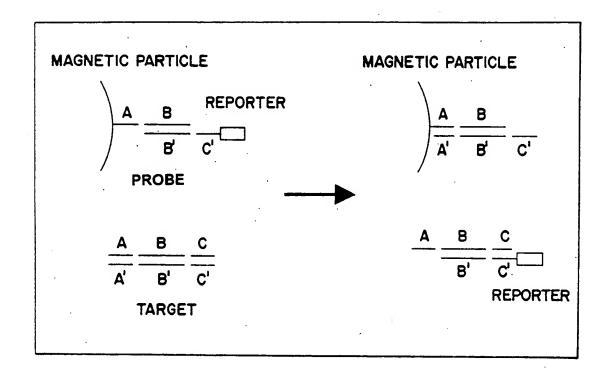
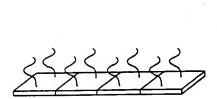
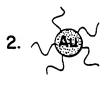


FIG. 25A 1. ∞ (TARGET)





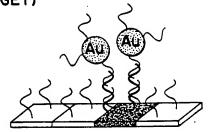
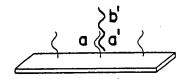
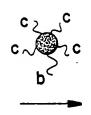


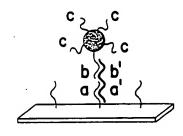
FIG. 25B

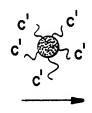


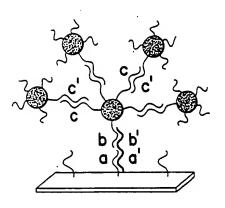














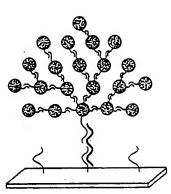


FIG. 26A

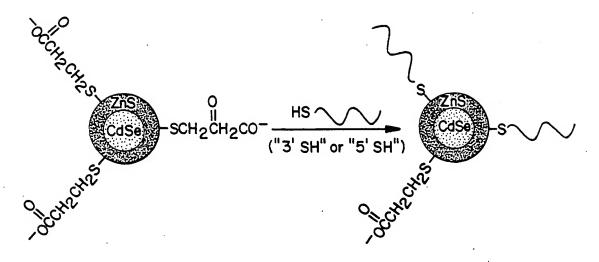
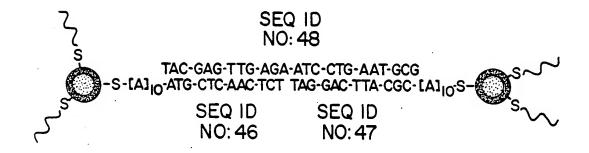


FIG. 26B



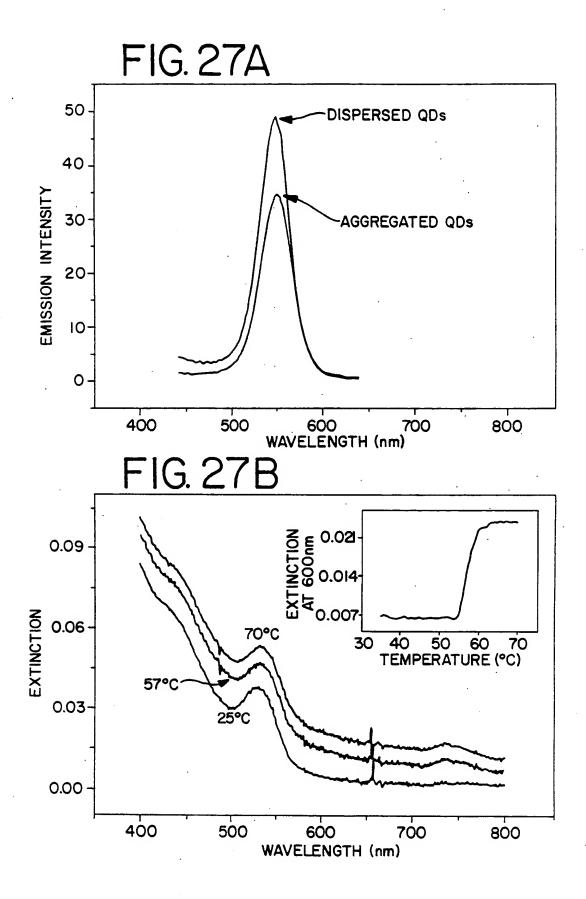


FIG. 27C

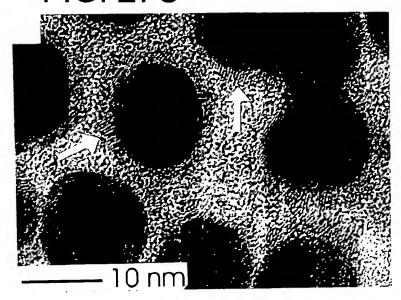


FIG. 27D

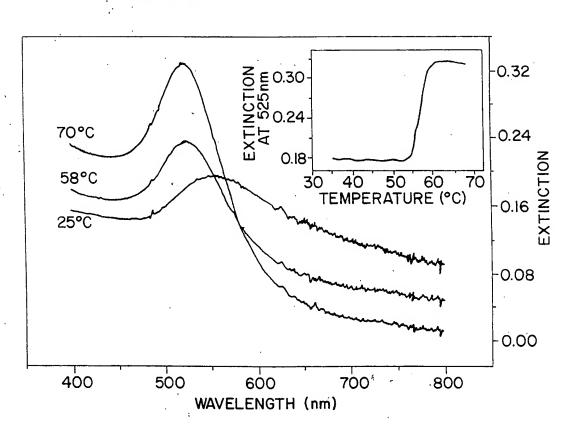


FIG. 28A

FIG. 28B

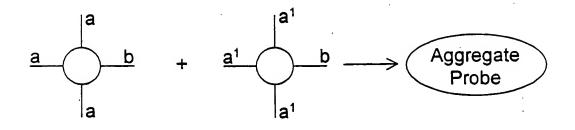


FIG. 28C

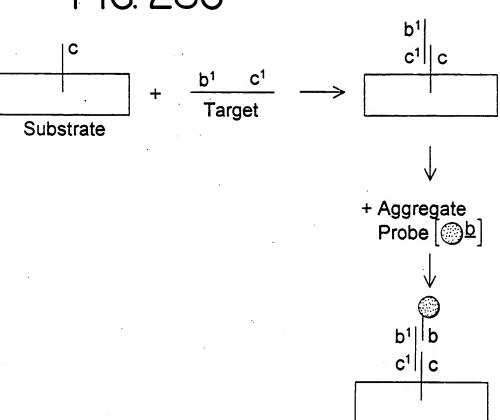


FIG. 28D

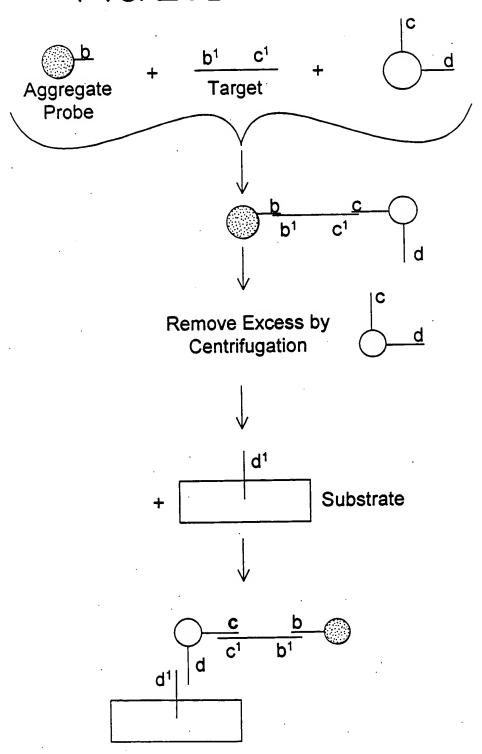
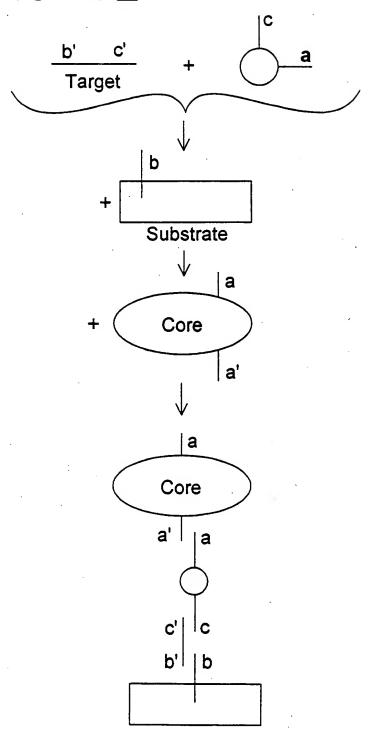
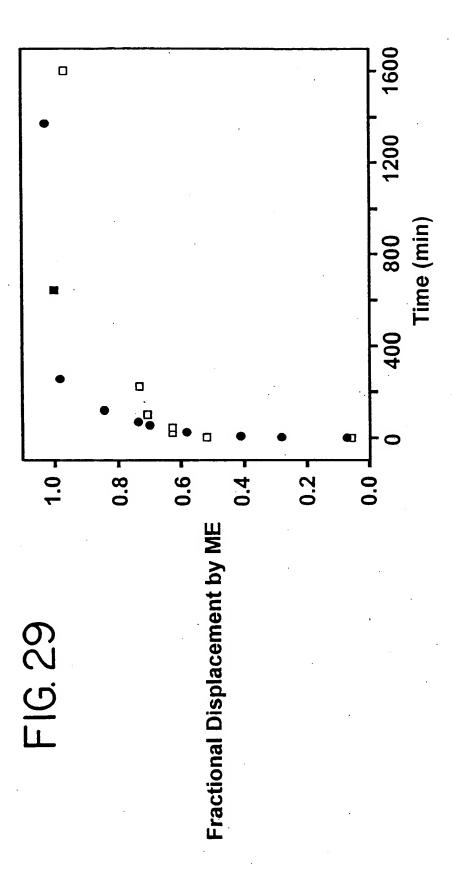
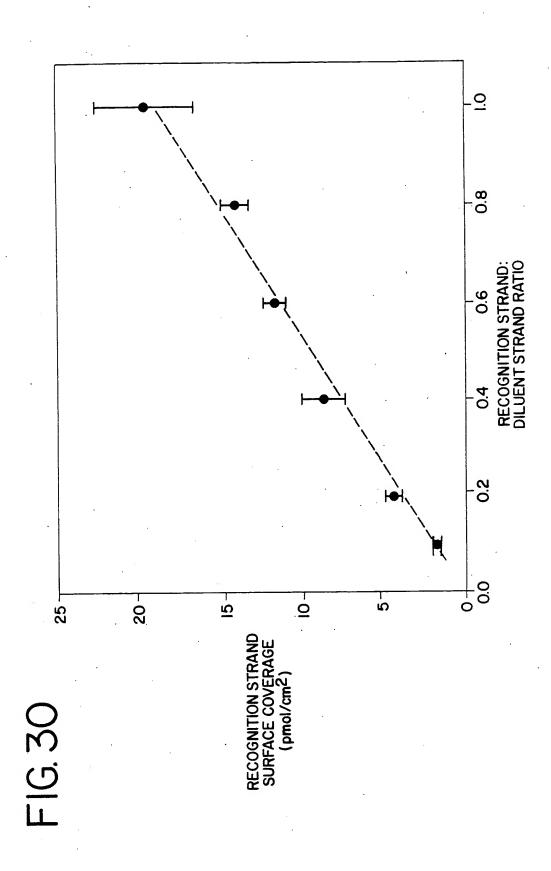


FIG. 28E







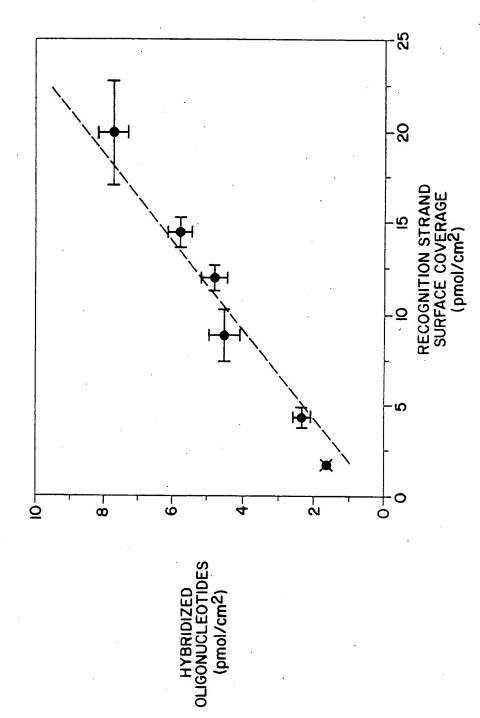
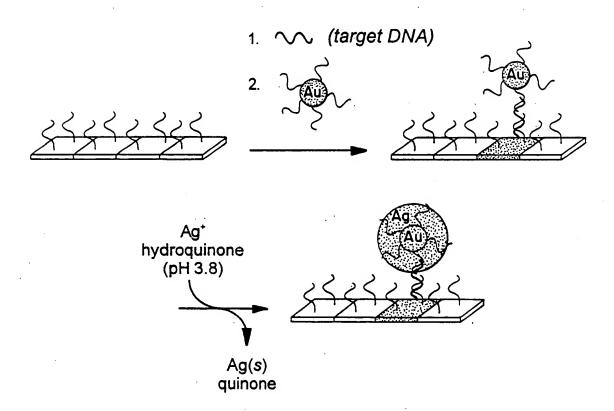


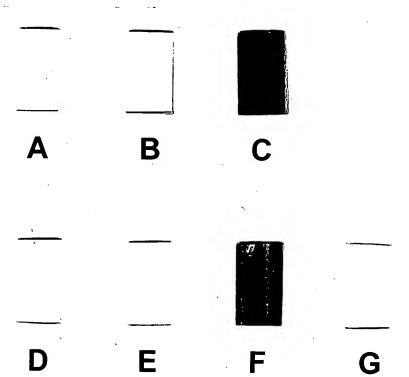
FIG. 31

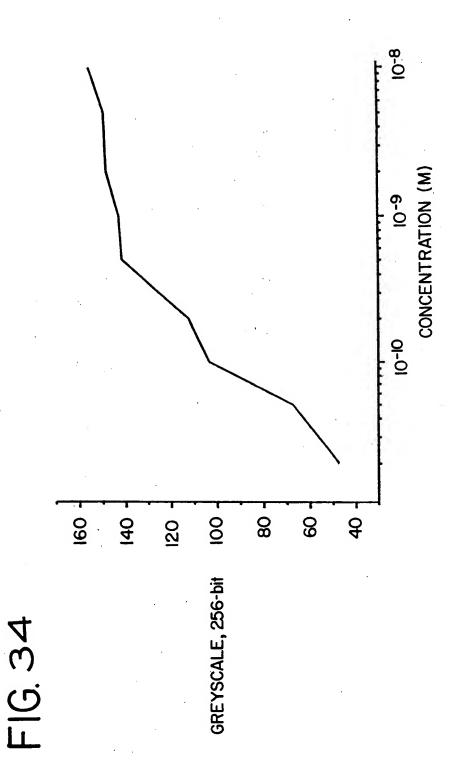
[SEQIDNO:56]

5' GGA T**T**A TTG TTA--AAT ATT GAT AAG GAT 3' CCT A**N**T AAC AAT TTA TAA CTA TTC CTA [SEQ ID NO: 57] [SEQ ID NO: 58]

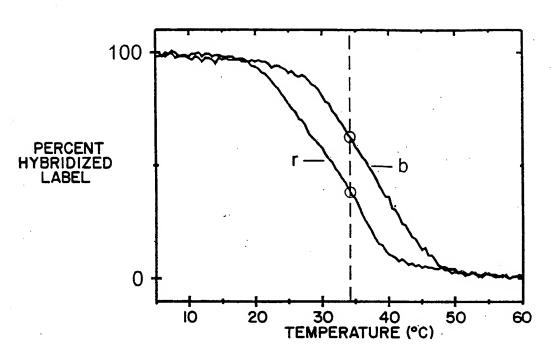
> N = A (complementary), G,C,T (mismatched)

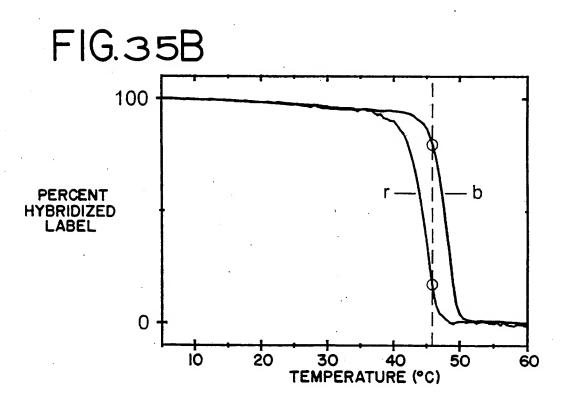












Replacement Sheet USSN 10/716,829 Mirkin et al.

FIG. 36A

FIG. 36B

C A T G

FIG.37A

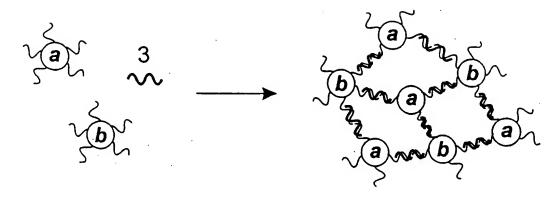
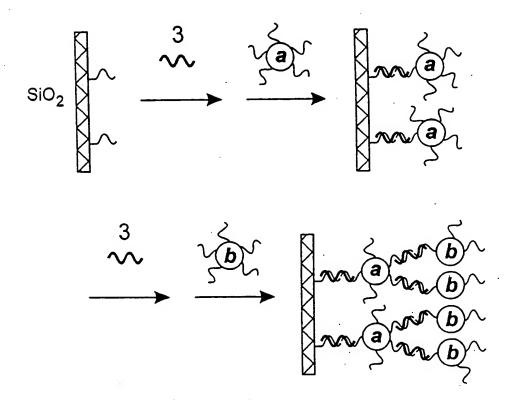
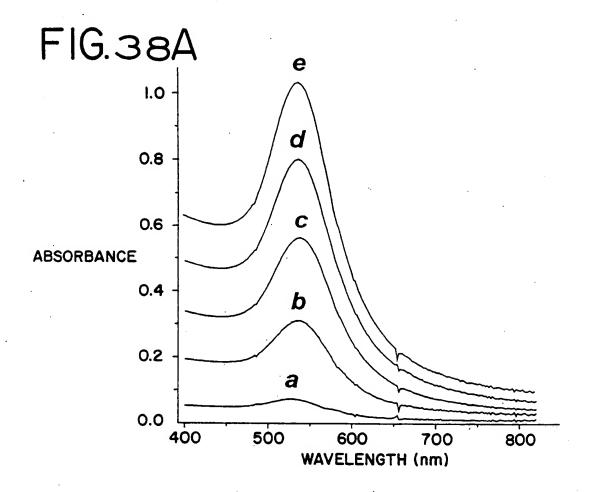


FIG.37B





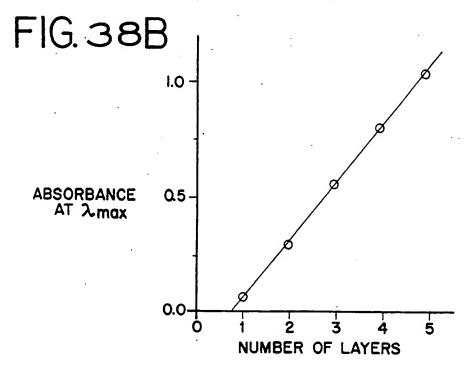


FIG. 39A

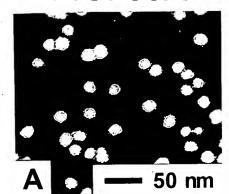
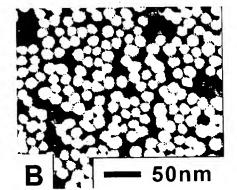
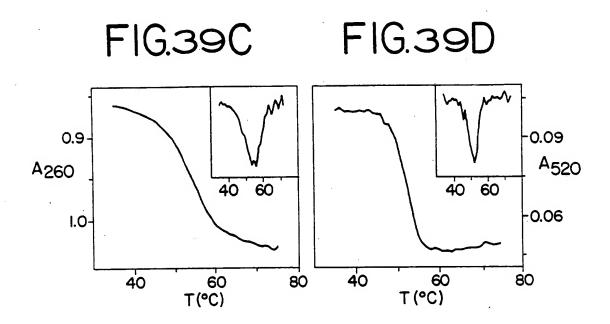
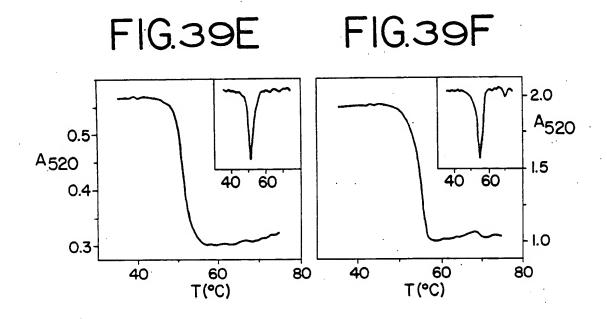


FIG. 39B







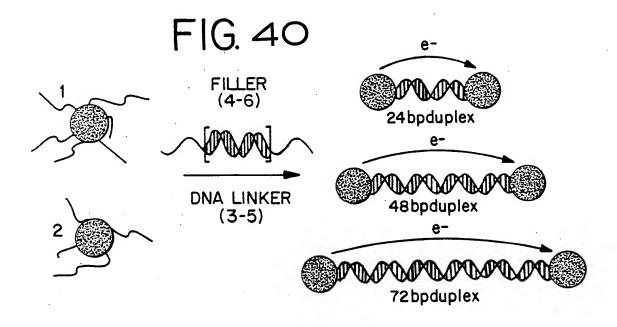
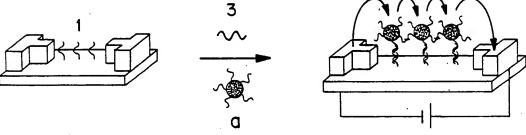


FIG. 41



II HS-(CH₂)₆OR₁

III $\overset{S-(CH_2)_6OR_1}{S-(CH_2)_6OH}$

 R_1

a = H $b = (iPr)_2NP(OCH_2CH_2CN)$ $c1 = 5'p(A_{20})$ -TATCGTTCCATCAGCT [SEQ ID NO: 65]

 $c2 = 5'-p(A_{20})$ -TTGATCTTCCGTTCT [SEQ ID NO: 66]

Target $\hat{I} = 79$ -mer oligonucleotide with target region:

3'-....ATAGCAAGGTAGTCGAGCAACTAGAAAGGCAAGA......5'
[SEQ ID NO: 67]

R₃ = hydrogen, an alkyl group, an aryl group, or a substituted alkyl or aryl group

 R_4 = an attached oligonucleotide or modified oligonucleotide